

FIG. 1

IND-1

Bis (2-ethylhexyl) phthalate
MW 390.6

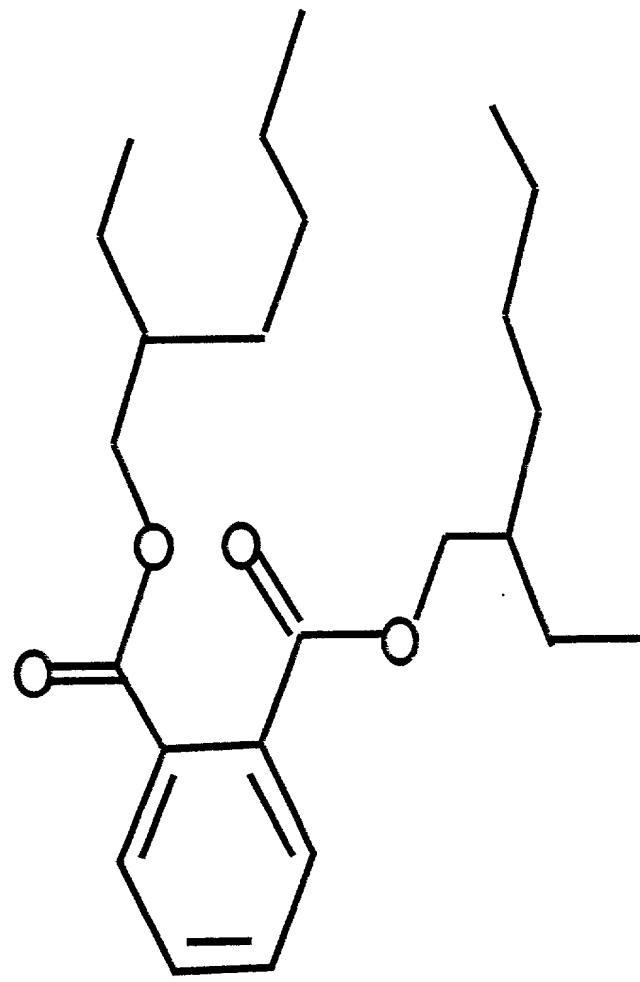
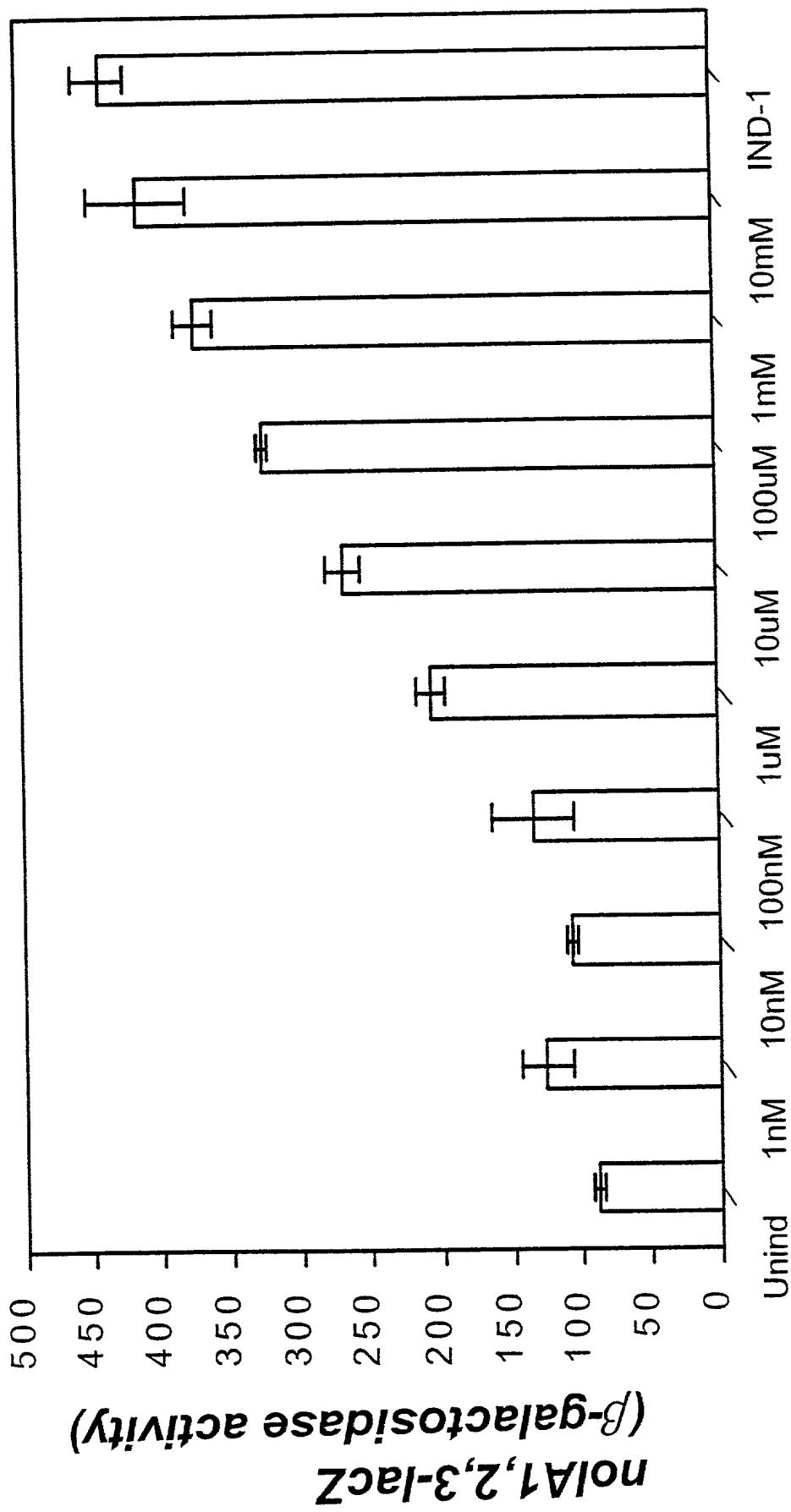


FIG. 2A



IND-1 (concentration)

FIG. 2B

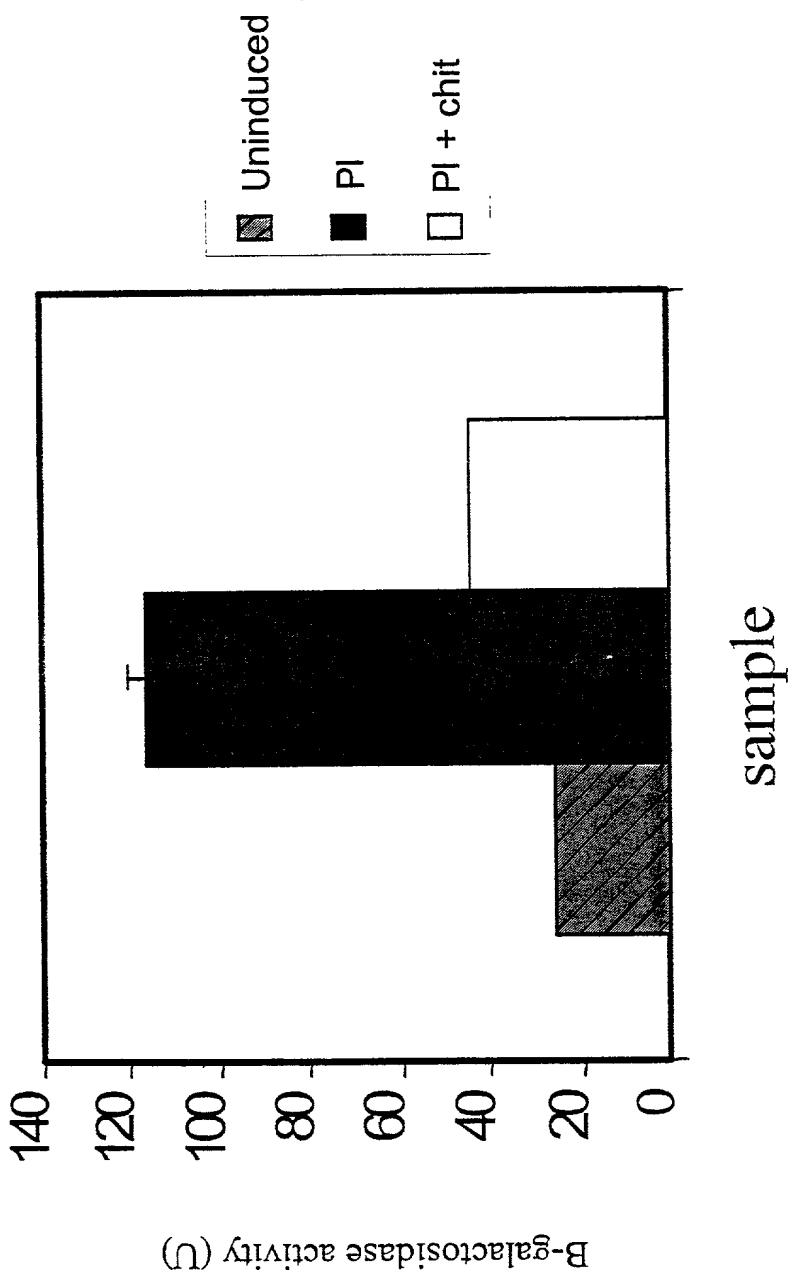


FIG. 3

sample

FIG. 4A

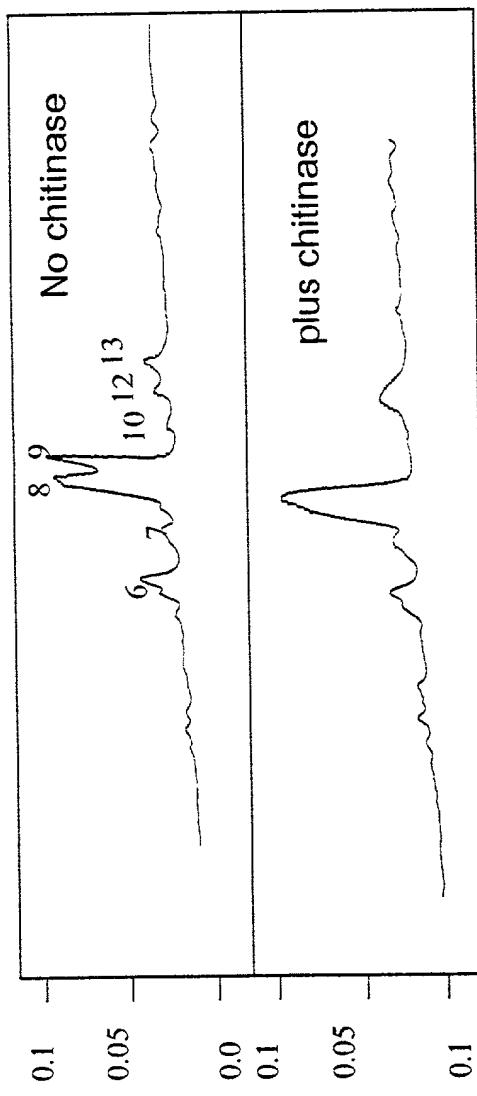


FIG. 4B



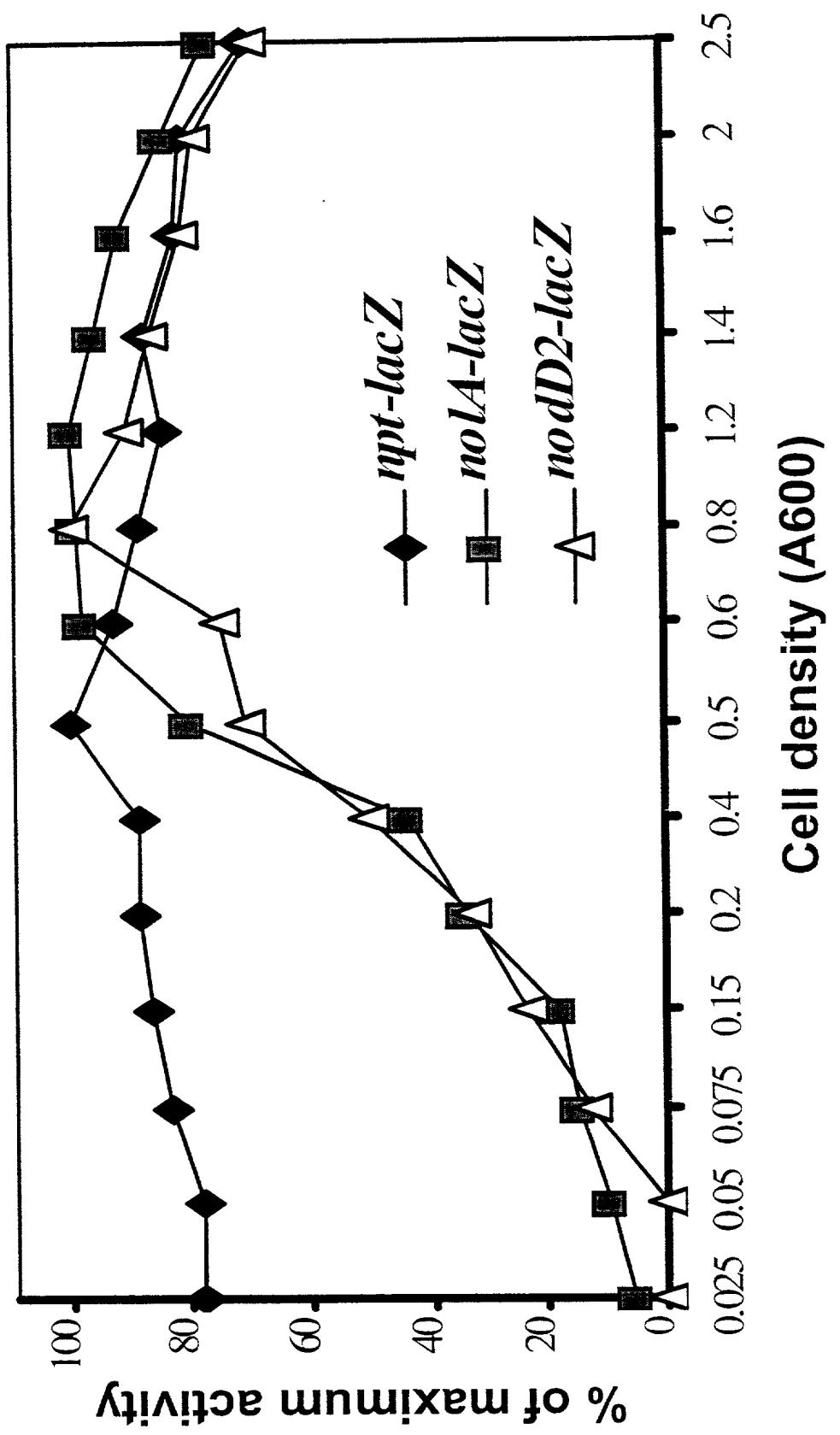


FIG. 5A

A₆₀₀ of culture at time of inducer extraction

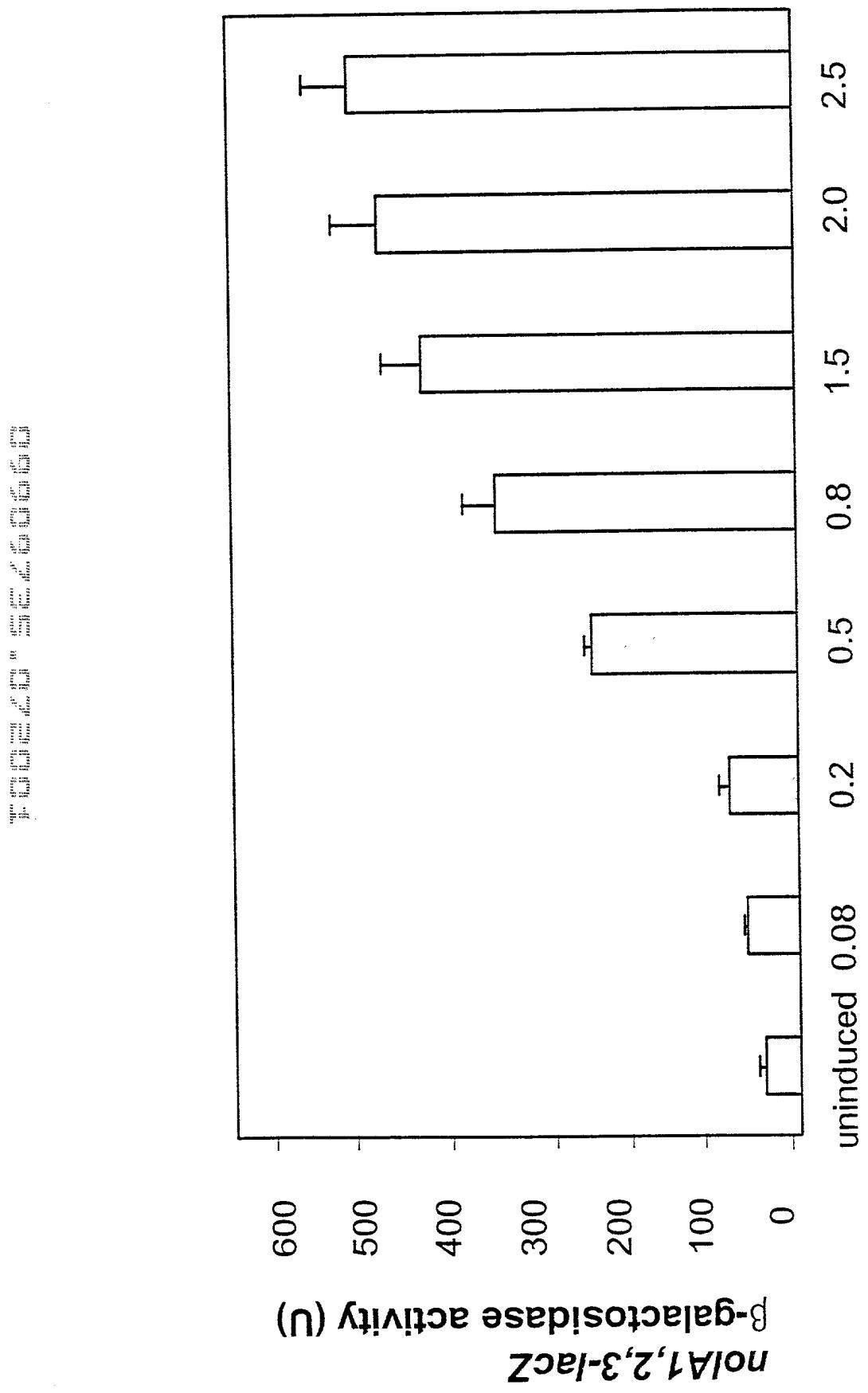


FIG. 5B

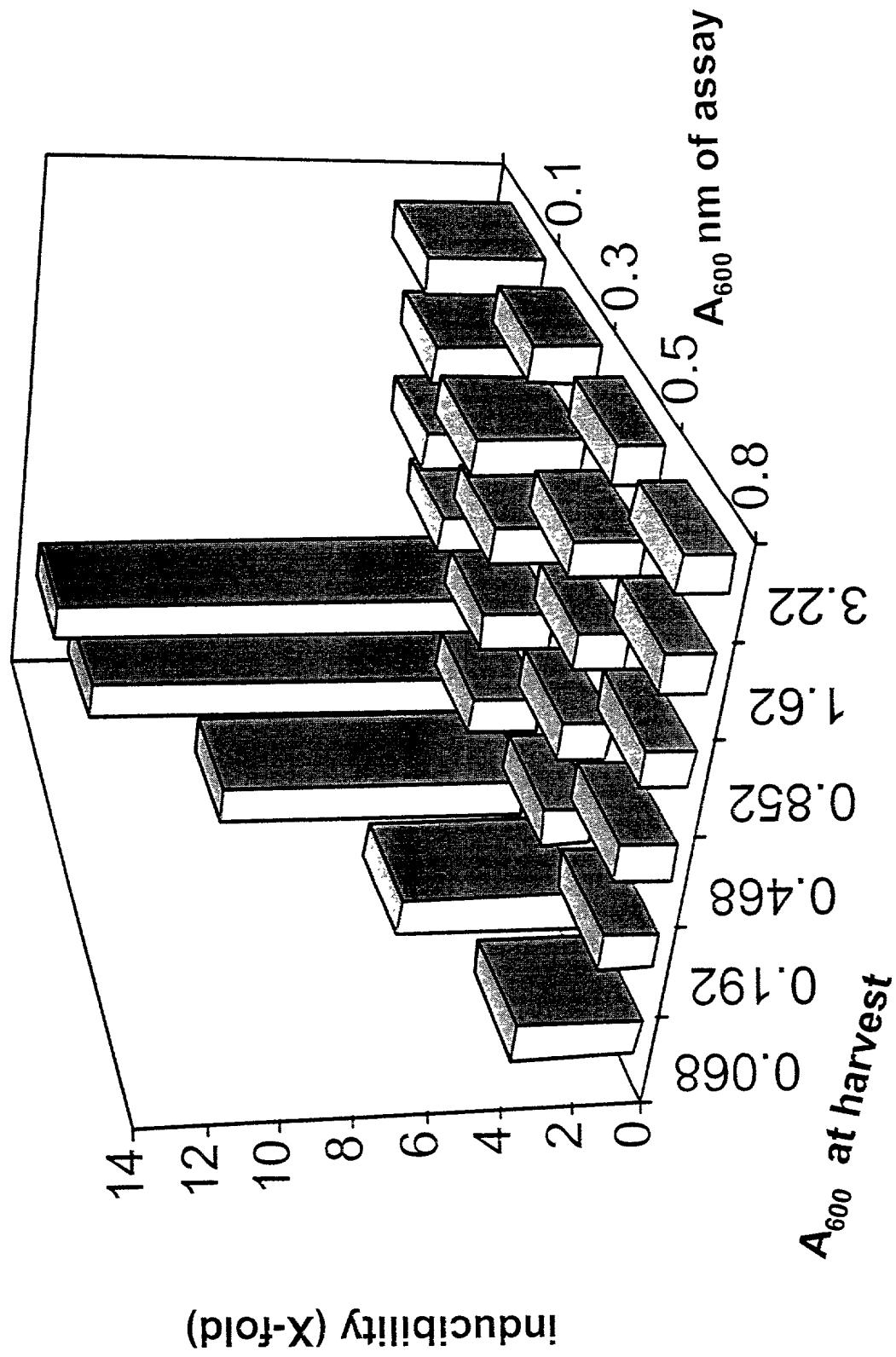
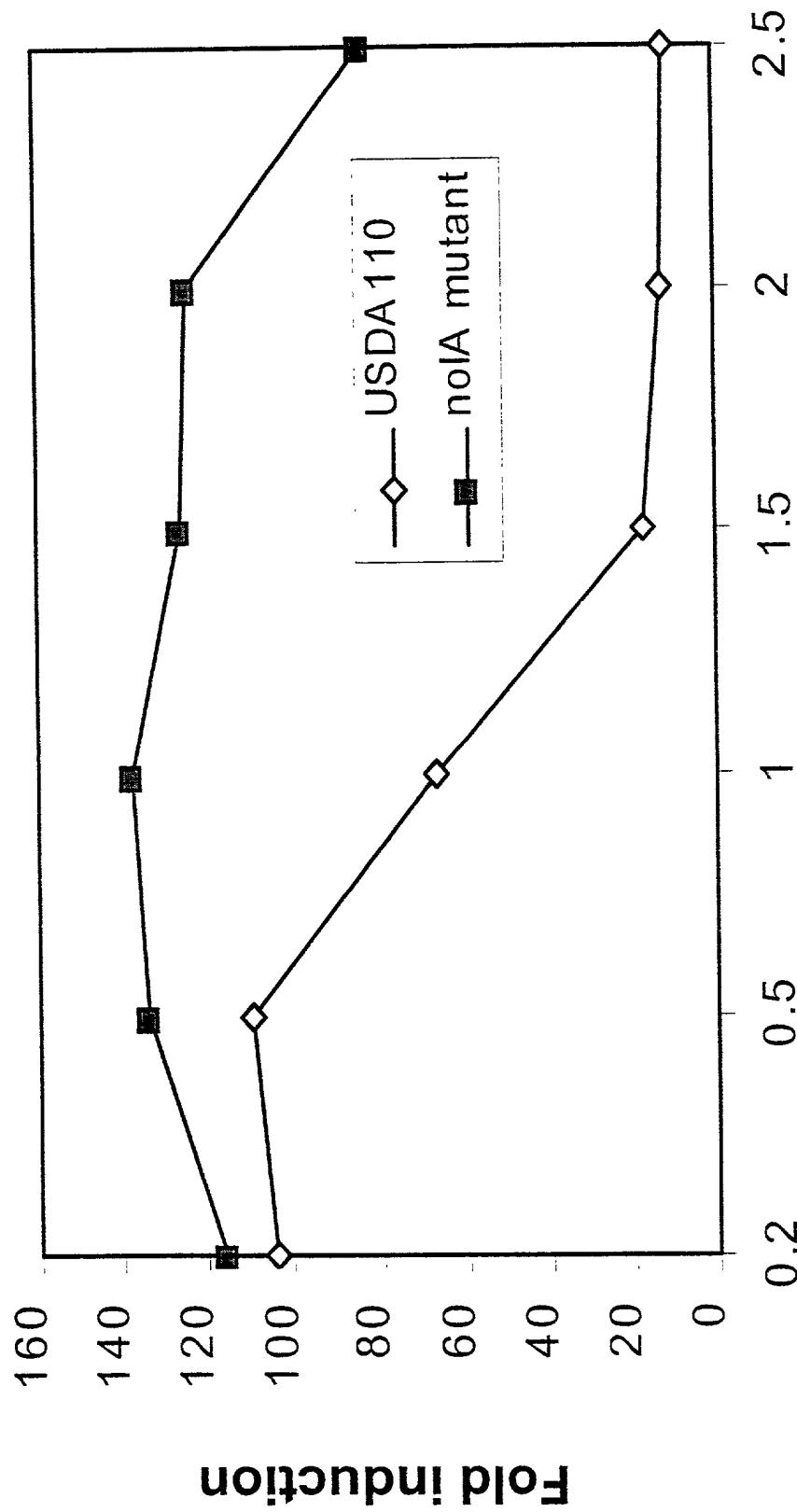


FIG. 6

Cell density (Δ 600)

FIG. 7



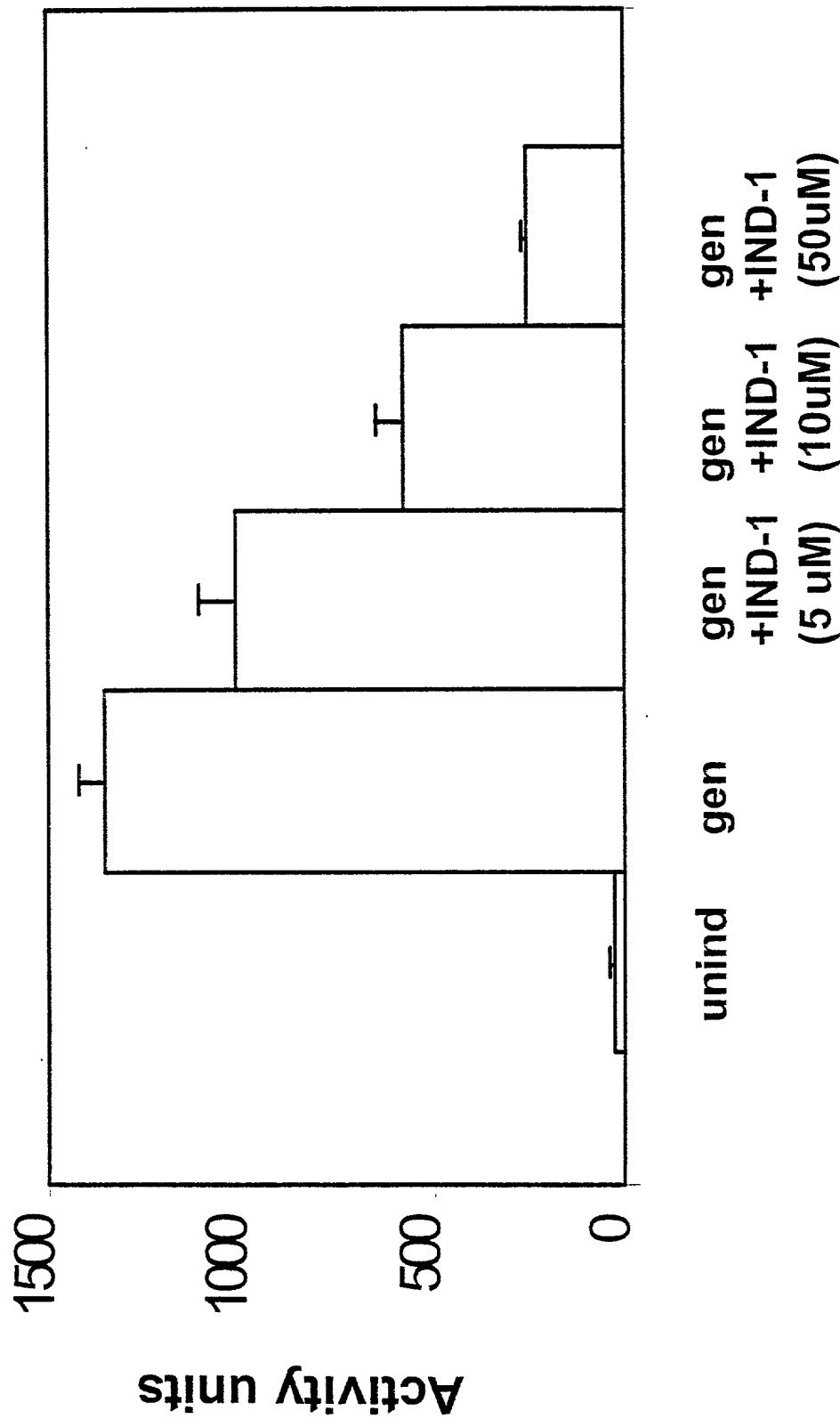
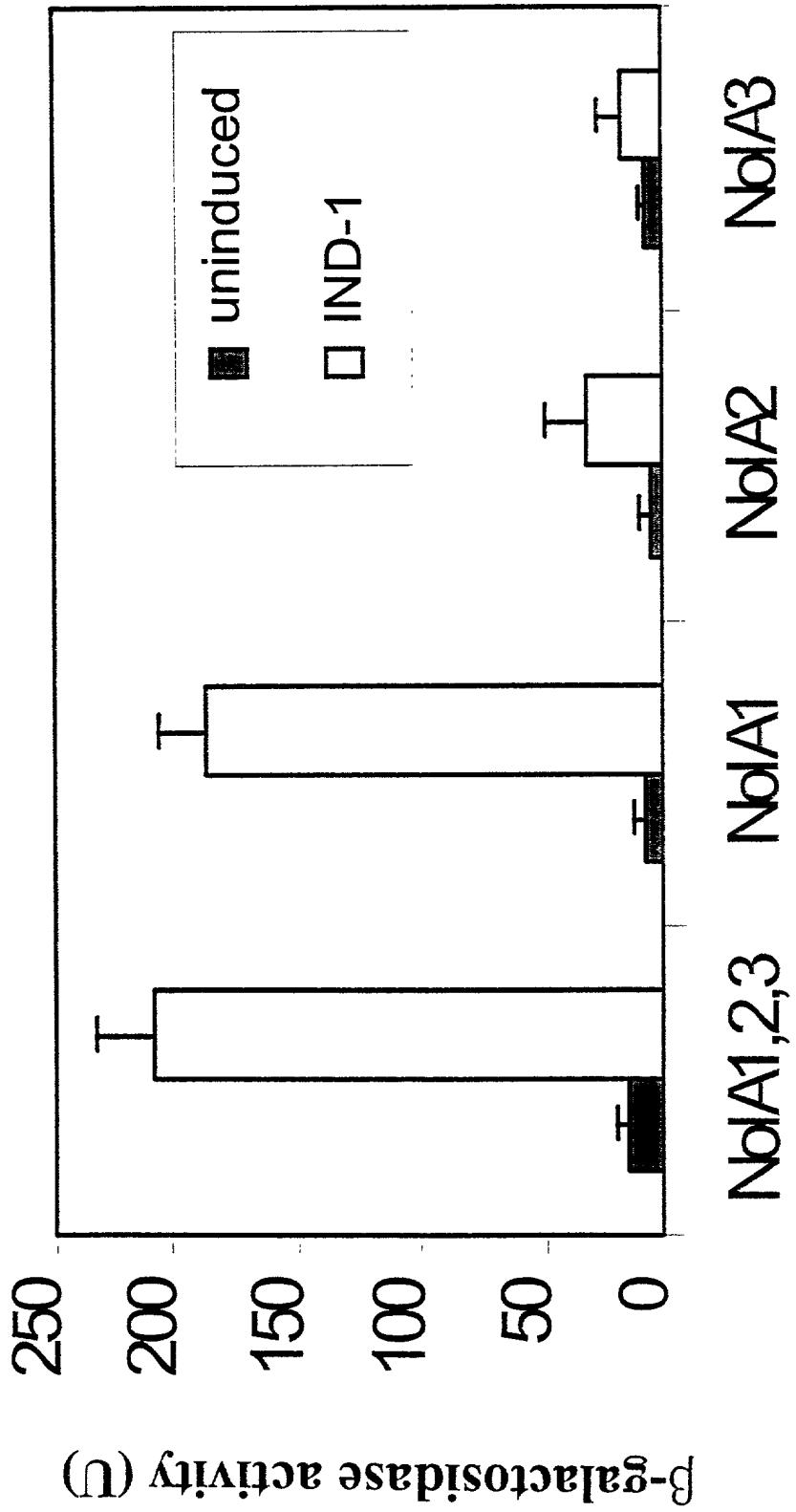


FIG. 8

FIG. 9

nolA-lacZ fusion



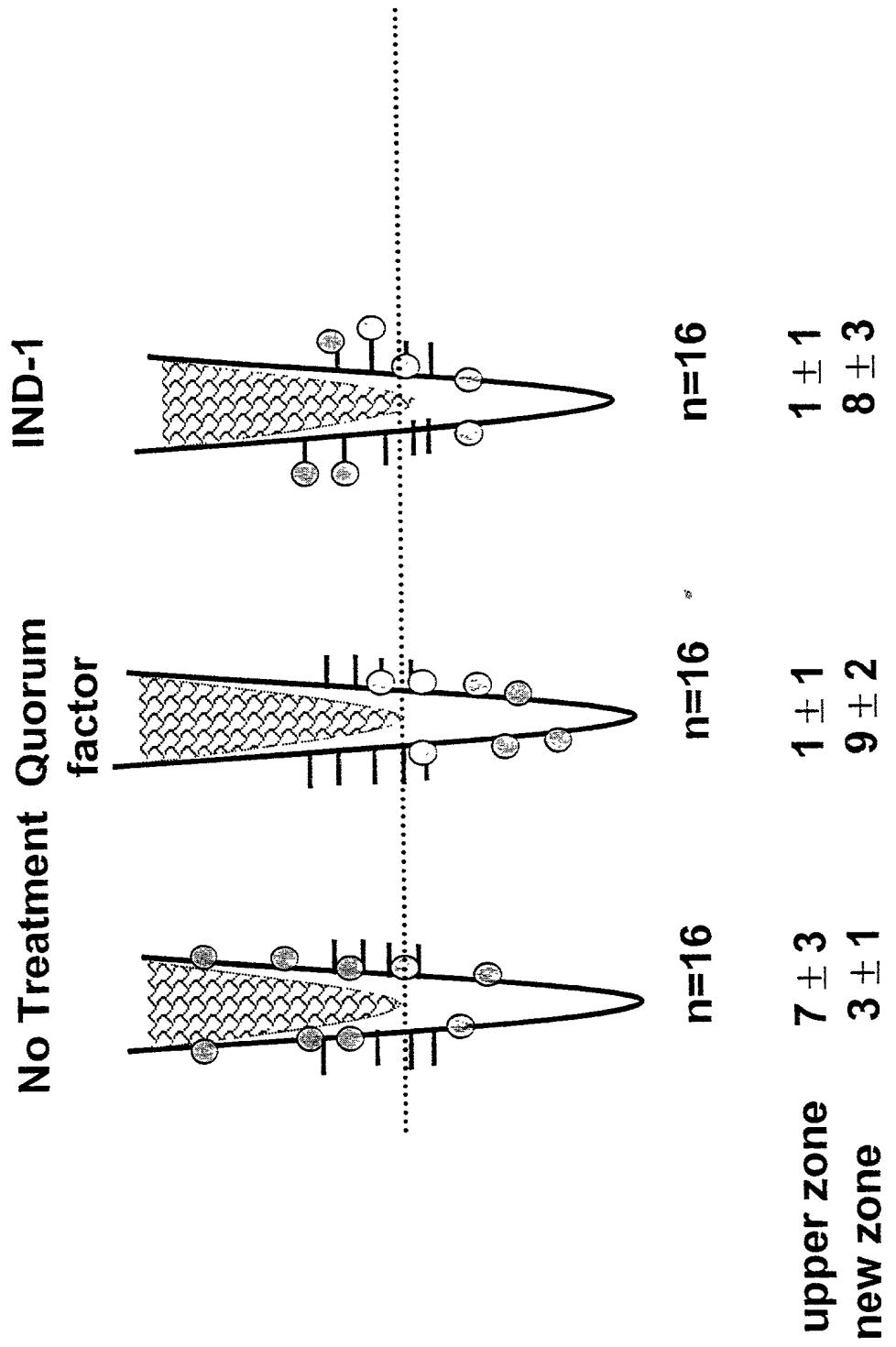
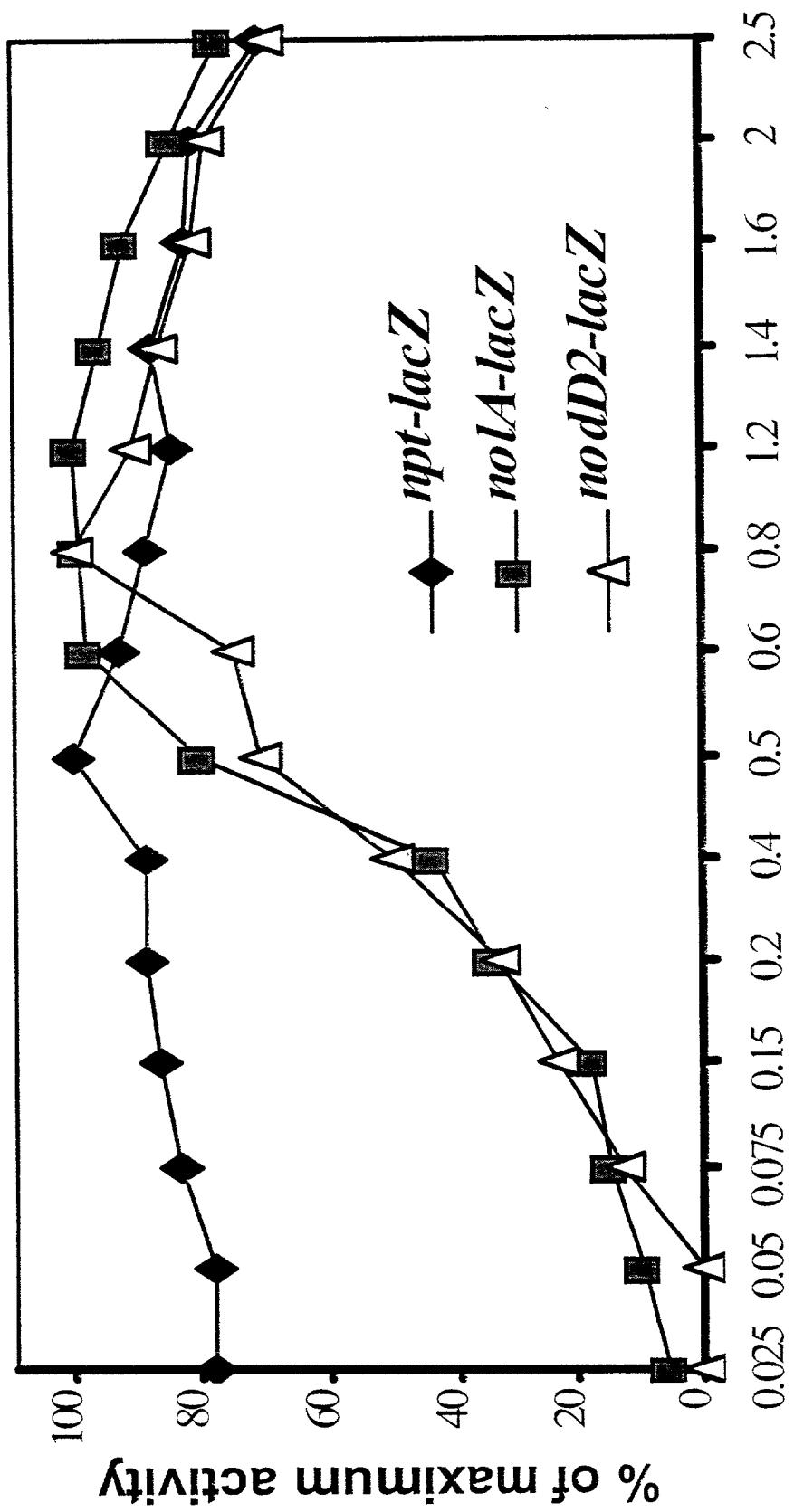


FIG. 10

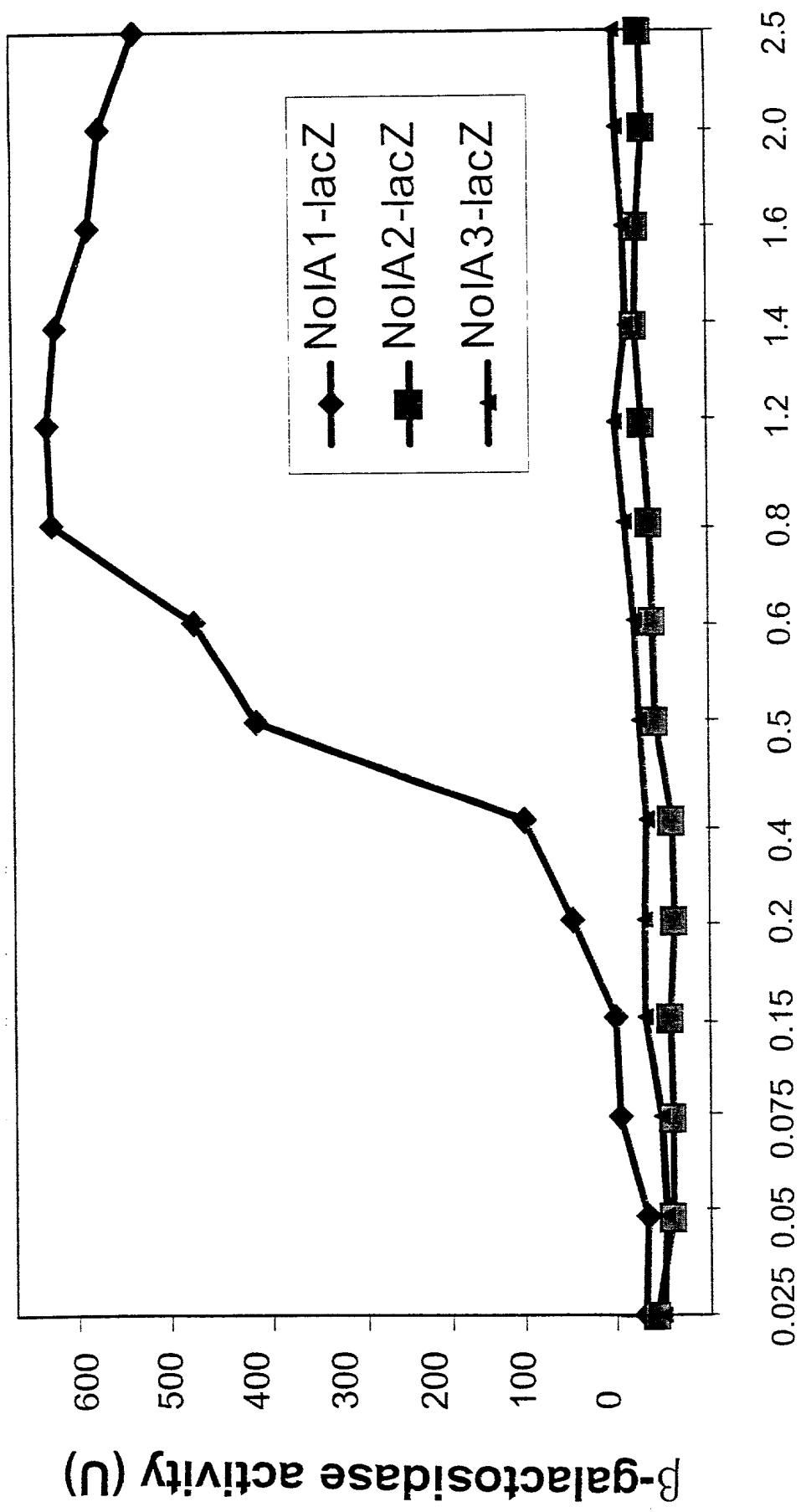
Cell density (A600)

FIG. 11A



Cell density (A_{600})

FIG. 11B



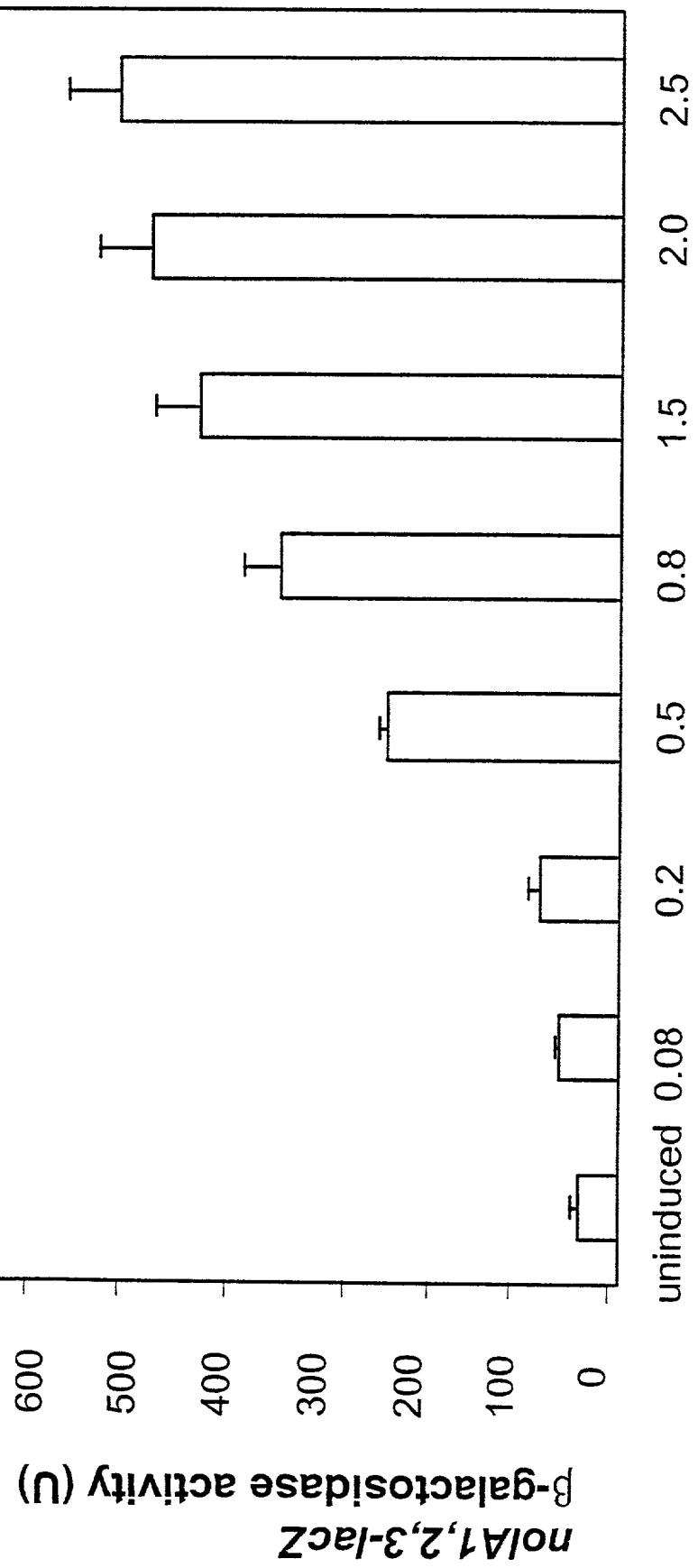


FIG. 11C

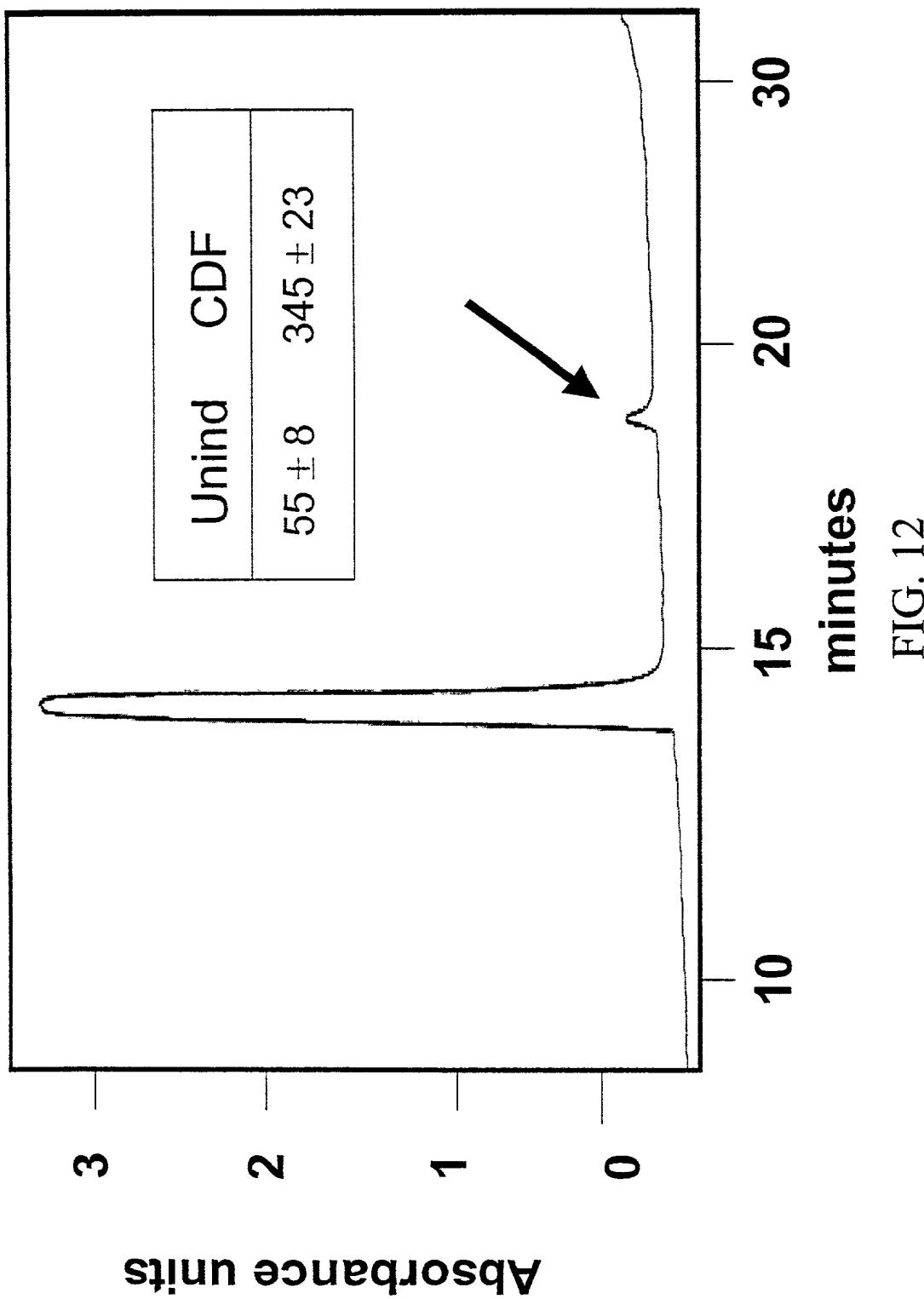
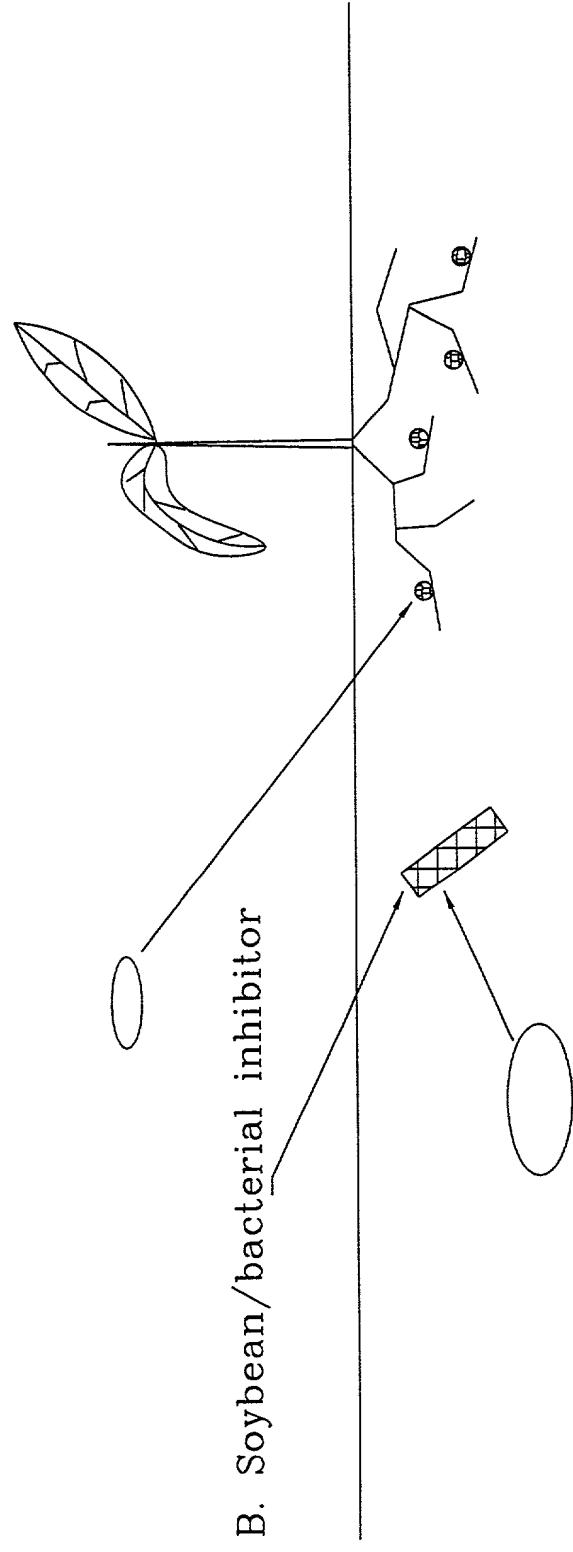


FIG. 12

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Inoculant

A. Inhibitor resistant *B. japonicum* (e.g., *Nola mutant*)



B. Soybean/bacterial inhibitor

Indigenous *B. japonicum* (sensitive to inhibitor)

FIG. 13

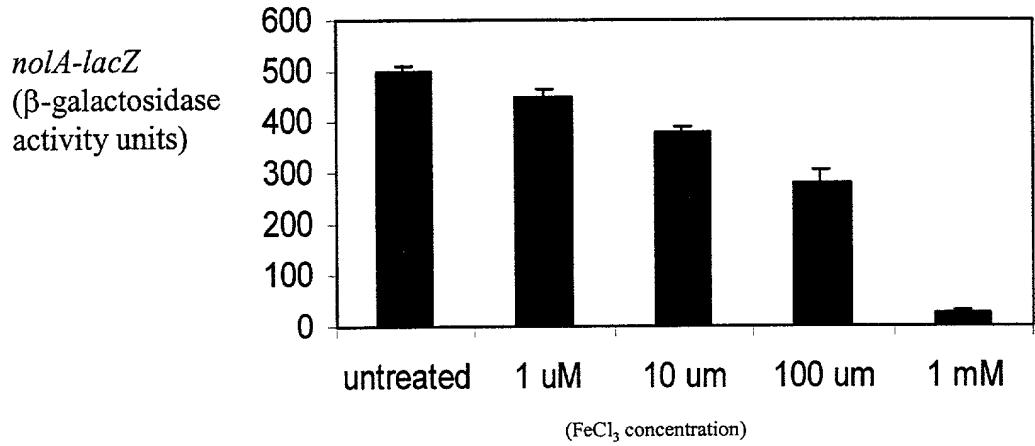


Figure 14A

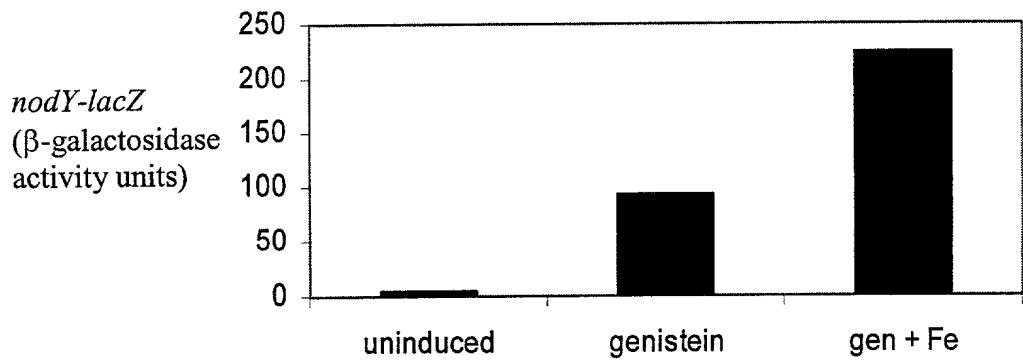


Figure 14B

RATIO OF NWSB:110	% OCCUPANCY BY NWSB MUTANT	
	Untreated	BEHP
1:10 (A)	0	11
(B)	2	0
10:1 (A)	83	95
(B)	93	92
1:1 (A)	57	78
1:1 (B)	40	74

Figure 15

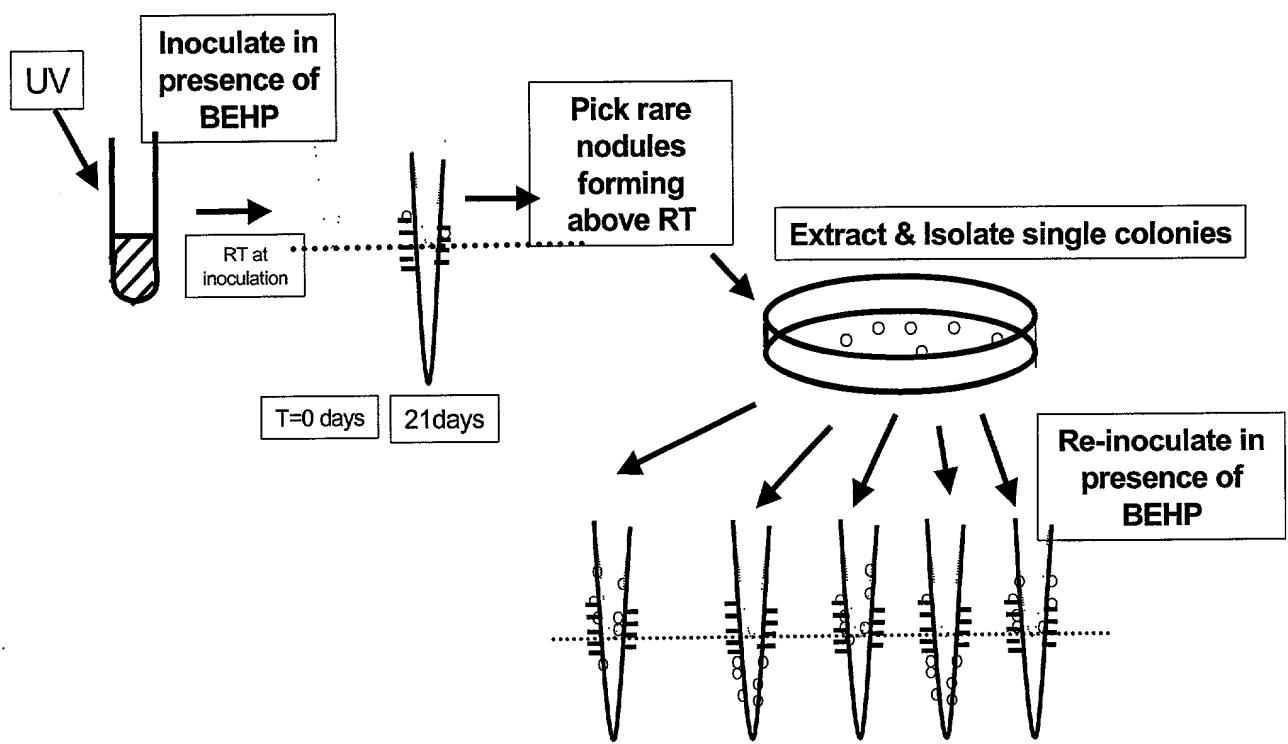


Figure 16

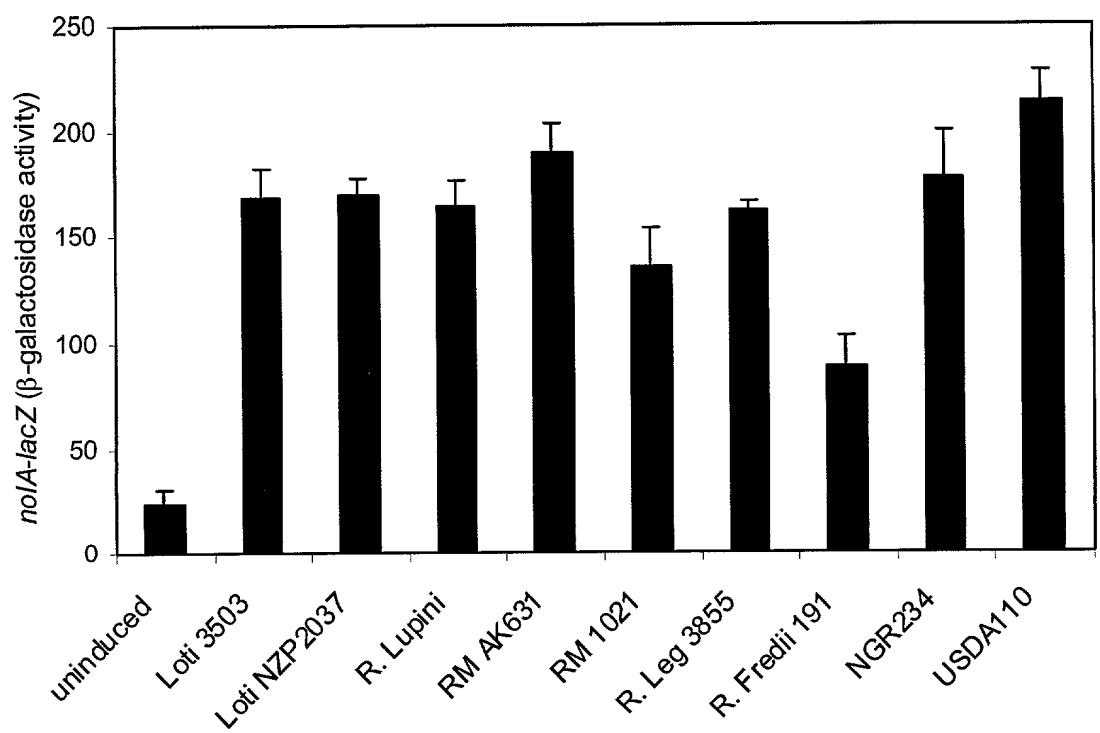


Figure 17

STRAIN	<i>NolA-lacZ</i> expression (fold induction)	+/- induction
<i>Bradyrhizobium japonicum</i> USDA 110	8.0	+++++
<i>Rhizobium Loti</i> NZP2037	3.0	++
<i>Rhizobium lupini</i>	2.8	++
<i>Sinorhizobium meliloti</i> AK631	3.5	++
<i>Sinorhizobium meliloti</i> 1021	2.4	+
<i>Rhizobium leguminosarum</i>	3.0	++
<i>Sinorhizobium</i> sp. NGR234	3.0	++
<i>Pseudomonas fluorescens</i> 5R	0.9	-
<i>Pseudomonas fluorescens</i> DFC50	0.8	-
<i>Pseudomonas aeruginosa</i> PAO1	1.0	-
<i>Pseudomonas syringae</i> B3A	1.1	-
<i>Pseudomonas syringae</i> B457	1.2	-
<i>Pseudomonas aureofaciens</i> Q2A7	1.0	-
<i>Agrobacterium</i> GV101	2.7	++
<i>Agrobacterium</i> LB4404	2.4	+
Marine isolate, gamma proteobacterium (Uwo.Ps)	1.2	-
Marine isolate, gamma proteobacterium (uwo.stk)	1.1	-
Marine isolate, gamma proteobacterium (uwo.mor)	0.9	-
<i>Aeromonas caviae</i>	1.8	-
<i>Vibrio harveyii</i>	2.4	-
<i>Vibrio natriegens</i>	1.3	-
<i>Vibrio splendidus</i>	2.5	-
<i>Rhodopseudomonas palustris</i>	2.7	++
<i>Salmonella typhi</i>	1.1	-
<i>Salmonella enteruditis</i>	1.0	-
<i>Salmonella typhi</i> 284	1.0	-
<i>M. smeraglitis</i>	1.0	-

FIG. 18